

02-03.ST25.txt  
SEQUENCE LISTING

<110> Quay, Steven C.  
<120> Compositions And Methods For Modulating Physiology Of Epithelial  
Junctional Adhesion Molecules For Enhanced Mucosal Delivery Of  
Therapeutic Compounds  
<130> 02-03US  
<150> 60/392,512  
<151> 2002-06-28  
<160> 900  
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Gly Val Asn Pro Thr Ala Gln Ser Ser  
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Gly Ser Leu Tyr Gly Ser Gln Ile Tyr  
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Ala Leu Cys Asn Gln Phe Tyr Thr Pro  
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Tyr Leu Tyr His Tyr Cys Val Val Asp  
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Pro Ala Ala Thr Gly Leu Tyr Val Asp Gln Tyr  
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Gln Phe Tyr Thr Pro Ala Ala Thr Gly Leu Tyr Val Asp Gln Tyr Leu  
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Tyr His Tyr

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Tyr Leu Tyr His Tyr Cys Val Val Asp  
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Gln Leu Tyr His Tyr Cys Val Val Asp Pro  
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Asp Gln Tyr Leu Tyr His Tyr Cys Val Val Asp Pro Gln  
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Val Asp Gln Tyr Leu Tyr His Tyr Cys Val Val Asp Pro Gln Glu

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Glu Ala Ile

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Gly Leu Tyr Val  
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Gly Ile Leu Arg Asp Phe Tyr Ser Pro Leu  
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Gly Phe Ser Leu Gly Leu Trp Met Glu Cys  
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Gly Ile Leu Arg Asp Phe Tyr Ser Pro Leu Val Pro Asp Ser  
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Arg Asp Phe Tyr Ser  
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Arg Asp Phe Tyr Ser  
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Arg Asp Phe Tyr  
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His Asn Ile Ile Gln Asp Phe Tyr Asn Pro Leu Val  
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Ala His Asn Ile Ile Gln Asp Phe Tyr Asn Pro Leu Val Ala  
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Thr Ala His Asn Ile Ile Gln Asp Phe Tyr Asn Pro Leu Val Ala Ser  
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Trp Thr Ala His Asn Ile Ile Gln Asp Phe Tyr Asn Pro Leu Val Ala  
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Ala Ser Gly Gln  
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&lt;400&gt; 81

Val Ser Trp Thr Ala His Asn Ile Ile Gln Asp Phe Tyr Asn Pro Leu  
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Val Ala Ser Gly Gln Lys  
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Pro Val Val Pro Glu Ala Gln Lys Arg Glu Met Gly  
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&lt;400&gt; 83

Asn Pro Val Val Pro Glu Ala Gln Lys Arg Glu Met Gly Ala  
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Phe Tyr Asn Pro Val Val Pro Glu Ala Gln Lys Arg Glu Met Gly Ala  
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Gly Leu

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Asp Phe Tyr Asn Pro Val Val  
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Arg Asp Phe Tyr Asn Pro Val Val Pro Glu Ala Gln Lys Arg Glu Met  
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Gly Ala Gly Leu Tyr Val  
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Lys Ile Thr Ala Ser Tyr Glu Asp Arg Val  
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Thr Phe Leu Pro Thr Gly Ile Thr Phe Lys  
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Ser Tyr Gly Glu Val Lys Val Lys Leu Ile  
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Val Leu Val Pro Pro Ser Lys Pro Thr Val  
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Tyr Gly Thr Pro Met Thr Ser Asn Ala Val  
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Arg Met Glu Ala Val Glu Arg Asn Val Gly  
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Ser Val Thr Val His Ser Ser Glu  
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Pro Glu Val Arg Ile Pro Glu Asn  
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Asn Pro Val Lys Leu Ser Cys Ala  
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Tyr Ser Gly Phe Ser Ser Pro Arg  
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Val Glu Trp Lys Phe Asp Gln Gly  
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Asp Thr Thr Arg Leu Val Cys Tyr  
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Asn Asn Lys Ile Thr Ala Ser Tyr  
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Glu Asp Arg Val Thr Phe Leu Pro  
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Thr Gly Ile Phe Lys Ser Val  
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Thr Arg Glu Asp Thr Gly Thr Tyr  
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Thr Cys Met Val Ser Glu Glu Gly  
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Gly Asn Ser Tyr Gly Glu Val Lys  
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Val Lys Leu Ile Val Leu Val Pro  
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Pro Ser Lys Pro Thr Val Asn Ile  
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Pro Ser Ser Ala Thr Ile Gly Asn  
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Arg Ala Val Leu Thr Cys Ser Glu  
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Gln Asp Gly Ser Pro Pro Ser Glu  
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Tyr Thr Trp Phe Lys Asp Gly Ile  
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Val Met Pro Thr Asn Pro Lys Ser  
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Thr Arg Ala Phe Ser Asn Ser Ser  
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Tyr Val Leu Asn Pro Thr Thr Gly  
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Glu Leu Val Phe Asp Pro Leu Ser  
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Ala Ser Asp Thr Gly Glu Tyr Ser  
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Cys Glu Ala Arg Asn Gly Tyr Gly  
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Arg Met Glu Ala Val Glu Arg Asn  
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Val Gly Val Ile  
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Ser Val Thr Val His  
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Ser Ser Glu Pro Glu Val Arg Ile Pro Glu  
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Asn Asn Pro Val Lys Leu Ser Cys Ala Tyr  
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Ser Gly Phe Ser Ser Pro Arg Val Glu Trp  
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Lys Phe Asp Gln Gly Asp Thr Thr Arg Leu  
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Val Cys Tyr Asn Asn Lys Ile Thr Ala Ser  
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Tyr Glu Asp Arg Val Thr Phe Leu Pro Thr  
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Gly Ile Thr Phe Lys Ser Val Thr Arg Glu  
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Asp Thr Gly Thr Tyr Thr Cys Met Val Ser  
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Glu Glu Gly Gly Asn Ser Tyr Gly Glu Val  
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Lys Val Lys Leu Ile Val Leu Val Pro Pro  
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Ser Lys Pro Thr Val Asn Ile Pro Ser Ser  
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<400> 149

Ala Thr Ile Gly Asn Arg Ala Val Leu Thr  
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Cys Ser Glu Gln Asp Gly Ser Pro Pro Ser  
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<400> 151

Glu Tyr Thr Trp Phe Lys Asp Gly Ile Val  
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Met Pro Thr Asn Pro Lys Ser Thr Arg Ala  
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Phe Ser Asn Ser Ser Tyr Val Leu Asn Pro  
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Thr Thr Gly Glu Leu Val Phe Asp Pro Leu  
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Ser Ala Ser Asp Thr Gly Glu Tyr Ser Cys  
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Glu Ala Arg Asn Gly Tyr Gly Thr Pro Met  
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Thr Ser Asn Ala Val Arg Met Glu Ala Val  
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Glu Arg Asn Val Gly Val Ile  
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Ser Val Thr Val  
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His Ser Ser Glu Pro Glu Val Arg  
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Ile Pro Glu Asn Asn Pro Val Lys  
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Leu Ser Cys Ala Tyr Ser Gly Phe  
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Ser Ser Pro Arg Val Glu Trp Lys  
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Phe Asp Gln Gly Asp Thr Thr Arg  
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Leu Val Cys Tyr Asn Asn Lys Ile  
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Thr Ala Ser Tyr Glu Asp Arg Val  
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Thr Phe Leu Pro Thr Gly Ile Thr  
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Phe Lys Ser Val Thr Arg Glu Asp  
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Thr Gly Thr Tyr Thr Cys Met Val  
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Ser Glu Glu Gly Gly Asn Ser Tyr  
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Gly Glu Val Lys Val Lys Leu Ile  
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Val Leu Pro Pro Ser Lys Pro  
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Thr Val Asn Ile Pro Ser Ser Ala  
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Thr Ile Gly Asn Arg Ala Val Leu  
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Thr Cys Ser Glu Gln Asp Gly Ser  
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Pro Pro Ser Glu Tyr Thr Trp Phe  
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Lys Asp Gly Ile Val Met Pro Thr  
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Asn Pro Lys Ser Thr Arg Ala Phe  
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Pro Thr Thr Gly Glu Leu Val Phe  
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Asp Pro Leu Ser Ala Ser Asp Thr  
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Gly Glu Tyr Ser Cys Glu Ala Arg  
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Asn Gly Tyr Gly Thr Pro Met Thr  
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Ser Asn Ala Val Arg Met Glu Ala  
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Val Glu Arg Asn Val Gly Val Ile  
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Pro Val Arg Ile Pro Glu  
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Pro Glu Val Arg Ile Pro Glu Asn  
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Ser Glu Pro Glu Val Arg Ile Pro Glu Asn Asn Pro  
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Ser Ser Glu Pro Glu Val Arg Ile Pro Glu Asn Asn Pro Val  
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His Ser Ser Glu Pro Glu Val Arg Ile Pro Glu Asn Asn Pro Val Lys  
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Val His Ser Ser Glu Pro Glu Val Arg Ile  
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Pro Glu Asn Asn Pro Val Lys Leu  
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Thr Val His Ser Ser Glu Pro Glu Val Arg Ile  
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Pro Glu Asn Asn Pro Val Lys Leu Ser  
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Val Arg Ile Pro Glu  
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Val Arg Ile Pro Glu Asn  
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Val Arg Ile Pro Glu Asn Asn  
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Val Arg Ile Pro Glu Asn Asn Pro Val  
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Val Arg Ile Pro Glu Asn Asn Pro Val Lys  
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Val Arg Ile Pro Glu Asn Asn Pro Val Lys Leu  
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Val Arg Ile Pro Glu Asn Asn Pro Val Lys Leu Ser  
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Glu Val Arg Ile Pro  
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Pro Glu Val Arg Ile Pro  
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Glu Pro Glu Val Arg Ile Pro  
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Ser Glu Pro Glu Val Arg Ile Pro  
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Ser Ser Glu Pro Glu Val Arg Ile Pro  
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Val Arg Val Pro  
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Pro Val Arg Val Pro Glu  
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Pro Glu Val Arg Val Pro Glu Asn  
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Glu Pro Glu Val Arg Val Pro Glu Asn Asn  
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Ser Glu Pro Glu Val Arg Val Pro Glu Asn Asn Pro  
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Ser Ser Glu Pro Glu Val Arg Val Pro Glu Asn Asn Pro Val  
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His Ser Ser Glu Pro Glu Val Arg Val Pro Glu Asn Asn Pro Val Lys  
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Val His Ser Ser Glu Pro Glu Val Arg Val Pro Glu Asn Asn Pro Val  
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Lys Leu

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Pro Glu Asn Asn Pro Val Lys Leu Ser  
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Val Arg Val Pro Glu  
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Val Arg Val Pro Glu Asn  
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Val Arg Val Pro Glu Asn Asn  
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Val Arg Val Pro Glu Asn Asn Pro Val  
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Val Arg Val Pro Glu Asn Asn Pro Val Lys  
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Val Arg Val Pro Glu Asn Asn Pro Val Lys Leu  
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Glu Val Arg Val Pro  
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Pro Glu Val Arg Val Pro  
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Phe Ser Ser Pro Arg  
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Arg Glu Asp Thr Gly Thr Tyr Thr Cys Met Val  
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Ser Val Thr Arg Glu Asp Thr Gly Thr Tyr Thr Cys Met  
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Lys Ser Val Thr Arg Glu Asp Thr Gly Thr Tyr Thr Cys Met  
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Glu Asp Thr Gly Thr Tyr Thr Cys Met Val Ser Glu Glu Gly  
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Val Thr Arg Glu Asp Thr Gly Thr Tyr Thr Cys Glu Val Ser Glu  
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Ser Val Thr Arg Glu Asp Thr Gly Thr Tyr Thr Cys Glu Val Ser Glu  
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Glu

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Glu Glu Gly

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Glu Asp Thr Gly Thr Tyr Thr Cys Glu Val Ser Glu Glu  
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&lt;400&gt; 428

Glu Asp Thr Gly Thr Tyr Thr Cys Glu Val Ser Glu Glu Gly  
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&lt;400&gt; 429

Glu Asp Thr Gly Thr Tyr Arg Cys Met  
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&lt;210&gt; 430

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&lt;400&gt; 430

Arg Glu Asp Thr Gly Thr Tyr Arg Cys Met Val  
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&lt;211&gt; 13

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

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&lt;223&gt; Synthetic construct

&lt;400&gt; 431

Thr Arg Glu Asp Thr Gly Thr Tyr Arg Cys Met Val Ser  
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Val Thr Arg Glu Asp Thr Gly Thr Tyr Arg Cys Met Val Ser Glu  
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Glu

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Glu Glu Gly

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<400> 435

Arg Glu Asp Thr Gly Thr Tyr Arg Cys Met  
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Glu Asp Thr Gly Thr Tyr Arg Cys Met Val Ser Glu Glu  
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Glu

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Glu Asp Thr Gly Thr Tyr Arg Cys Glu Val  
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Glu Asp Thr Gly Thr Tyr Arg Cys Glu Val Ser  
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Glu Asp Thr Gly Thr Tyr Arg Cys Glu Val ser Glu  
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Glu Asp Thr Gly Thr Tyr Arg Cys Glu Val ser Glu Glu  
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Arg Glu Asp Ser Gly Thr Tyr Thr Cys Met Val  
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Val Thr Arg Glu Asp Ser Gly Thr Tyr Thr Cys Met Val Ser Glu  
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Glu Asp Ser Gly Thr Tyr Thr Cys Glu Val Ser Glu Glu Gly  
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Pro Val Val Gln Glu Phe Glu Ser Val Glu  
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Leu Ser Cys Ile Ile Thr Asp Ser Gln Thr  
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Asp Asn Lys Ile Gln Gly Asp Leu Ala Gly  
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Pro Val Gly Lys Met Ala Thr Leu His Cys  
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Arg Asn Ser Ser Phe His Leu Asn Ser Glu  
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Thr Gly Thr Leu Val Phe Thr Ala Val His  
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Lys Asp Asp Ser Gly Gln Tyr Tyr Cys Ile  
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Ala Val Asn Leu Lys  
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Ser Ser Asn Arg Thr Pro Val Val Gln Glu  
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Glu Trp Lys Lys Ile Gln Asp Glu Gln Thr  
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Thr Tyr Val Phe Phe Asp Asn Lys Ile Gln  
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Ile Asp Glu Ile Val Ile Glu Leu Thr Val  
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Ala Thr Leu His Cys Gln Glu Ser Glu Gly  
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His Leu Asn Ser Glu Thr Gly Thr Leu Val  
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Phe Thr Ala Val His Lys Asp Asp Ser Gly  
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Gln Tyr Tyr Cys Ile Ala Ser Asn Asp Ala  
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Gly Ser Ala Arg Cys Glu Glu Gln Glu Met  
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Ile Thr Asp Ser Gln Thr Ser Asp  
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Pro Arg Ile Glu Trp Lys Lys Ile  
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Gln Asp Glu Gln Thr Thr Tyr Val  
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Phe Phe Asp Asn Lys Ile Gln Gly  
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Asp Leu Ala Gly Arg Ala Glu Ile  
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Leu Gly Lys Thr Ser Leu Lys Ile  
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Trp Asn Val Thr Arg Arg Asp Ser  
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Ala Leu Tyr Arg Cys Glu Val Val  
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Ala Arg Asn Asp Arg Lys Glu Ile  
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Asp Glu Ile Val Ile Glu Leu Thr  
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Val Gln Val Lys Pro Val Thr Pro  
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Val Cys Arg Val Pro Lys Ala Val  
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Pro Val Gly Lys Met Ala Thr Leu  
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His Cys Gln Glu Ser Glu Gly His  
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Pro Arg Pro His Tyr Ser Trp Tyr  
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Arg Asn Asp Val Pro Leu Pro Thr  
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Asp Ser Arg Ala Asn Pro Arg Phe  
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Arg Asn Ser Ser Phe His Leu Asn  
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Ser Glu Thr Gly Thr Leu Val Phe  
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Thr Ala Val His Lys Asp Asp Ser  
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Gly Gln Tyr Tyr Cys Ile Ala Ser  
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Asn Asp Ala Gly Ser Ala Arg Cys  
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Glu Glu Gln Glu Met Glu Val Tyr  
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Asp Leu Asn  
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Ala Val Asn Leu  
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Lys Ser Ser Asn Arg Thr Pro Val  
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Val Gln Glu Phe Glu Ser Val Glu  
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Leu Ser Cys Ile Ile Thr Asp Ser  
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Gln Thr Ser Asp Pro Arg Ile Glu  
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Thr Thr Tyr Val Phe Phe Asp Asn  
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Arg Ala Glu Ile Leu Gly Lys Thr  
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Ser Leu Lys Ile Trp Asn Val Thr  
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Arg Arg Asp Ser Ala Leu Tyr Arg  
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Cys Glu Val Val Ala Arg Asn Asp  
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Arg Lys Glu Ile Asp Glu Ile Val  
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Ile Glu Leu Thr Val Gln Val Lys  
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Pro Val Thr Pro Val Cys Arg Val  
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Pro Lys Ala Val Pro Val Gly Lys  
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Met Ala Thr Leu His Cys Gln Glu  
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Ser Glu Gly His Pro Arg Pro His  
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Tyr Ser Trp Tyr Arg Asn Asp Val  
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Pro Leu Pro Thr Asp Ser Arg Ala  
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Asn Pro Arg Phe Arg Asn Ser Ser  
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Phe His Leu Asn Ser Glu Thr Gly  
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Thr Leu Val Phe Thr Ala Val His  
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Lys Asp Asp Ser Gly Gln Tyr Tyr  
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Cys Ile Ala Ser Asn Asp Ala Gly  
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Ser Ala Arg Cys Glu Glu Gln Glu  
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Met Glu Val Tyr Asp Leu Asn  
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Gly Phe Ser Ala Pro Lys Asp Gln Gln Val  
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Val Thr Ala Val Glu Tyr Gln Glu Ala Ile  
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Leu Ala Cys Lys Thr Pro Lys Lys Thr Val  
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Ser Ser Arg Leu Glu Trp Lys Lys Leu Gly  
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Arg Ser Val Ser Phe Val Tyr Tyr Gln Gln  
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Thr Leu Gln Gly Asp Phe Lys Asn Arg Ala  
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Glu Met Ile Asp Phe Asn Ile Arg Ile Lys  
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Asn Val Thr Arg Ser Asp Ala Gly Lys Tyr  
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Arg Cys Glu Val Ser Ala Pro Ser Glu Gln  
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Gly Gln Asn Leu Glu Glu Asp Thr Val Thr  
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Leu Glu Val Leu Val Ala Pro Ala Val Pro  
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Ser Cys Glu Val Pro Ser Ser Ala Leu Ser  
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Gly Thr Val Val Glu Leu Arg Cys Gln Asp  
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Lys Glu Gly Asn Pro Ala Pro Glu Tyr Thr  
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Trp Phe Lys Asp Gly Ile Arg Leu Leu Glu  
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Asn Pro Arg Leu Gly Ser Gln Ser Thr Asn  
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Ser Ser Tyr Thr Met Asn Thr Lys Thr Gly  
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Thr Leu Gln Phe Asn Thr Val Ser Lys Leu  
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Asp Thr Gly Glu Tyr Ser Cys Glu Ala Arg  
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Asn Ser Val Gly Tyr Arg Arg Cys Pro Gly  
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Lys Arg Met Gln Val Asp Asp Leu Asn  
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Gly Phe Ser Ala Pro  
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Lys Asp Gln Gln Val Val Thr Ala Val Glu  
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Tyr Gln Glu Ala Ile Leu Ala Cys Lys Thr  
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Pro Lys Lys Thr Val Ser Ser Arg Leu Glu  
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Trp Lys Lys Leu Gly Arg Ser Val Ser Phe  
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Val Tyr Tyr Gln Gln Thr Leu Gln Gly Asp  
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Phe Lys Asn Arg Ala Glu Met Ile Asp Phe  
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Asn Ile Arg Ile Lys Asn Val Thr Arg Ser  
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Asp Ala Gly Lys Tyr Arg Cys Glu Val Ser  
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Ala Pro Ser Glu Gln Gly Gln Asn Leu Glu  
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Glu Asp Thr Val Thr Leu Glu Val Leu Val  
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Ala Pro Ala Val Pro Ser Cys Glu Val Pro  
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Ser Ser Ala Leu Ser Gly Thr Val Val Glu  
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Leu Arg Cys Gln Asp Lys Glu Gly Asn Pro  
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Ala Pro Glu Tyr Thr Trp Phe Lys Asp Gly  
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Ile Arg Leu Leu Glu Asn Pro Arg Leu Gly  
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Ser Gln Ser Thr Asn Ser Ser Tyr Thr Met  
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Asn Thr Lys Thr Gly Thr Leu Gln Phe Asn  
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Thr Val Ser Lys Leu Asp Thr Gly Glu Tyr  
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Ser Cys Glu Ala Arg Asn Ser Val Gly Tyr  
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Arg Arg Cys Pro Gly Lys Arg Met Gln Val  
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Asp Asp Leu Asn  
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Gly Phe Ser Ala Pro Lys Asp Gln  
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Gln Val Val Thr Ala Val Glu Tyr  
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Gln Glu Ala Ile Leu Ala Cys Lys  
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Thr Pro Lys Lys Thr Val Ser Ser  
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Arg Leu Glu Trp Lys Lys Leu Gly  
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Arg Ser Val Ser Phe Val Tyr Tyr  
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Gln Gln Thr Leu Gln Gly Asp Phe  
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Lys Asn Arg Ala Glu Met Ile Asp  
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Phe Asn Ile Arg Ile Lys Asn Val  
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Thr Arg Ser Asp Ala Gly Lys Tyr  
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Arg Cys Glu Val Ser Ala Pro Ser  
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Glu Gln Gly Gln Asn Leu Glu Glu  
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Asp Thr Val Thr Leu Glu Val Leu  
1 5

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Val Ala Pro Ala Val Pro Ser Cys  
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Glu Val Pro Ser Ser Ala Leu Ser  
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Gly Thr Val Val Glu Leu Arg Cys  
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Gln Asp Lys Glu Gly Asn Pro Ala  
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Pro Glu Tyr Thr Trp Phe Lys Asp  
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<400> 683

Gly Ile Arg Leu Leu Glu Asn Pro  
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Arg Leu Gly Ser Gln Ser Thr Asn  
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Ser Ser Tyr Thr Met Asn Thr Lys  
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Thr Gly Thr Leu Gln Phe Asn Thr  
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Val Ser Lys Leu Asp Thr Gly Glu  
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Val Gly Tyr Arg Arg Cys Pro Gly  
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Gly Phe Ser Ala  
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Pro Lys Asp Gln Gln Val Val Thr  
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Ala Val Glu Tyr Gln Glu Ala Ile  
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Leu Ala Cys Lys Thr Pro Lys Lys  
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Lys Lys Leu Gly Arg Ser Val Ser  
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Phe Val Tyr Tyr Gln Gln Thr Leu  
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Gln Gly Asp Phe Lys Asn Arg Ala  
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Glu Met Ile Asp Phe Asn Ile Arg  
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Ile Lys Asn Val Thr Arg Ser Asp  
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Ala Gly Lys Tyr Arg Cys Glu Val  
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Ser Ala Pro Ser Glu Gln Gly Gln  
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Asn Leu Glu Glu Asp Thr Val Thr  
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Val Pro Ser Cys Glu Val Pro Ser  
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Ser Ala Leu Ser Gly Thr Val Val  
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Glu Leu Arg Cys Gln Asp Lys Glu  
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Gly Asn Pro Ala Pro Glu Tyr Thr  
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Gln Ser Thr Asn Ser Ser Tyr Thr  
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Met Asn Thr Lys Thr Gly Thr Leu  
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Val Asp Asp Leu Asn  
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Tyr Ala Gly Asp Asn Ile Val Thr Ala Gln  
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Met Thr Pro Val Asn Ala Arg Tyr Glu Phe  
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Arg Ile Tyr Ser Tyr Ala Gly Asp Asn Ile  
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Val Thr Ala Gln Ala Met Tyr Glu Gly Leu  
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Trp Met Ser Cys Val Ser Gln Ser Thr Gly  
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Gln Ile Gln Cys Lys Val Phe Asp Ser Leu  
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Arg Ile Tyr Ser Tyr  
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Ala Gly Asp Asn Ile Val Thr Ala Gln Ala  
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Met Tyr Glu Gly Leu Trp Met Ser Cys Val  
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Ser Gln Ser Thr Gly Gln Ile Gln Cys Lys  
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Val Phe Asp Ser Leu Leu Asn Leu Ser Ser  
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Gln Glu Phe Tyr Asp Pro Met Thr  
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Pro Val Asn Ala Arg Tyr Glu  
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Ala Arg Tyr Glu  
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Lys Thr Ser Ser Tyr Val Gly Ala Ser Ile  
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Val Thr Ala Val Gly Phe Ser Lys Gly Leu  
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Trp Met Glu Cys Ala Thr His Ser Thr Gly  
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Ile Thr Gln Cys Asp Ile Tyr Ser Thr Leu  
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Leu Gly Leu Pro Ala Asp Ile Gln Ala Ala Gln  
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Lys Thr Ser Ser Tyr  
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Phe Ser Lys Gly Leu Trp Met Glu Cys Ala  
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Thr His Ser Thr Gly Ile Thr Gln Cys Asp  
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Asp Ile Gln Ala Ala Gln  
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Arg Asp Phe Tyr Ser Pro Leu  
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Val Pro Asp Ser Met Lys Phe Glu  
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Arg Val Thr Ala Phe Ile Gly Ser Asn Ile  
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Val Thr Ser Gln Thr Ile Trp Glu Gly Leu  
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Trp Met Asn Cys Val Val Gln Ser Thr Gly  
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Gln Met Gln Cys Lys Val Tyr Asp Ser Leu  
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Leu Ala Leu Pro Gln Asp Leu Gln Ala Ala Arg  
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Arg Val Thr Ala Phe  
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Ile Gly Ser Asn Ile Val Thr Ser Gln Thr  
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Val Tyr Asp Ser Leu Leu Ala Leu Pro Gln  
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Asp Leu Gln Ala Ala Arg  
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Gln Asp Phe Tyr Asn Pro Leu Val  
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Ala Ser Gly Gln Lys Arg Glu  
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Gln Val Thr Ala Phe Leu Asp His Asn Ile  
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Val Thr Ala Gln Thr Thr Trp Lys Gly Leu  
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His Met Gln Cys Lys Val Tyr Asp Ser Val  
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Gln Val Thr Ala Phe  
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Thr Trp Lys Gly Leu Trp Met Ser Cys Val  
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Val Gln Ser Thr Gly His Met Gln Cys Lys  
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Glu Val Gln Ala Ala Arg  
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Arg Glu Phe Tyr Asp Pro Ser Val  
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Pro Val Ser Gln Lys Tyr Glu  
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Arg Val Ser Ala Phe Ile Gly Ser Asn Ile  
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Ile Thr Ser Gln Asn Ile Trp Glu Gly Leu  
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Arg Val Ser Ala Phe  
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Arg Asp Phe Tyr Asn Pro Val Val  
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Pro Glu Ala Gln Lys Arg Glu  
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Lys Val Thr Ala Phe Ile Gly Asn Ser Ile  
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Val Val Ala Gln Val Val Trp Glu Gly Leu  
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Lys Val Thr Ala Phe  
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Ile Gly Asn Ser Ile Val Val Ala Gln Val  
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Val Trp Glu Gly Leu Trp Met Ser Cys Val  
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Arg Asp Phe Tyr Asn Pro Leu Val  
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Ala Glu Ala Gln Lys Arg Glu  
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Gln Met Ser Ser Tyr Ala Gly Asp Asn Ile  
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Ile Thr Ala Gln Ala Met Tyr Lys Gly Leu  
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Trp Met Asp Cys Val Thr Gln Ser Thr Gly  
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Met Met Ser Cys Lys Met Tyr Asp Ser Val  
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Leu Ala Leu Ser Ala Ala Leu Gln Ala Thr Arg  
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Gln Met Ser Ser Tyr  
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Ala Gly Asp Asn Ile Ile Thr Ala Gln Ala  
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Met Tyr Lys Gly Leu Trp Met Asp Cys Val  
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Thr Gln Ser Thr Gly Met Met Ser Cys Lys  
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Met Tyr Asp Ser Val Leu Ala Leu Ser Ala  
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Ala Leu Gln Ala Thr Arg  
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Thr Asp Phe Tyr Asn Pro Leu Ile  
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Pro Thr Asn Ile Lys Tyr Glu  
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Arg Val Ser Ala Phe Ile Glu Asn Asn Ile  
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Val Val Phe Glu Asn Phe Trp Glu Gly Leu  
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Trp Met Asn Cys Val Arg Gln Ala Asn Ile  
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Arg Met Gln Cys Lys Ile Tyr Asp Ser Leu  
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Leu Ala Leu Ser Pro Asp Leu Gln Ala Ala Arg  
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Arg Val Ser Ala Phe  
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<400> 804

Phe Trp Glu Gly Leu Trp Met Asn Cys Val  
 1 5 10

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<400> 805

Arg Gln Ala Asn Ile Arg Met Gln Cys Lys  
 1 5 10

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<400> 806

Ile Tyr Asp Ser Leu Leu Ala Leu Ser Pro  
 1 5 10

<210> 807  
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<400> 807

Arg Asp Phe Tyr Asn Ser Ile Val  
 1 5

<210> 808  
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<400> 808

Asn Val Ala Gln Lys Arg Glu  
 1 5

<210> 809  
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<400> 809

Lys Val Thr Ala Phe Ile Gly Asn Ser Ile  
 1 5 10

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<400> 810

Ala Glu Ala Leu Lys Arg Glu  
 1 5

<210> 811  
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<400> 811

Lys Val Ser Thr Ile Asp Gly Thr Val Ile  
 1 5 10

<210> 812  
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<400> 812

Thr Thr Ala Thr Tyr Trp Ala Asn Leu Trp  
 1 5 10

<210> 813  
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<400> 813

Lys Ala Cys Val Thr Asp Ser Thr Gly Val  
 1 5 10

<210> 814  
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<400> 814

Ser Asn Cys Lys Asp Phe Pro Ser Met Leu  
 1 5 10

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<400> 815

Ala Leu Asp Gly Tyr Ile Gln Ala Cys Arg  
1 5 10

<210> 816  
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<400> 816

Lys Val Ser Thr Ile  
1 5

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<400> 817

Asp Gly Thr Val Ile Thr Thr Ala Thr Tyr  
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<400> 818

Trp Ala Asn Leu Trp Lys Ala Cys Val Thr  
1 5 10

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<400> 819

Asp Ser Thr Gly Val Ser Asn Cys Lys Asp  
1 5 10

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<400> 820

Phe Pro Ser Met Leu Ala Leu Asp Gly Tyr  
 1 5 10

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<400> 821

Ile Gln Ala Cys Arg  
 1 5

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<400> 822

Glu Phe Phe Asp Pro Leu Phe  
 1 5

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<400> 823

Val Glu Gln Lys Tyr Glu  
 1 5

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<400> 824

Asp Arg Gly Tyr Gly Thr Ser Leu Leu Gly  
1 5 10

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<400> 825

Gly Ser Val Gly Tyr Pro Tyr Gly Gly Ser  
1 5 10

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<220>  
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<400> 826

Gly Phe Gly Ser Tyr Gly Ser Gly Tyr Gly  
1 5 10

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<400> 827

Tyr Gly Tyr Gly Tyr Gly Tyr Gly Tyr Gly  
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<400> 828

Gly Tyr Thr Asp Pro Arg  
1 5

<210> 829  
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<212> PRT  
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<220>  
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<400> 829

Asp Arg Gly Tyr Gly  
 1 5

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<220>  
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<400> 830

Thr Ser Leu Leu Gly Gly Ser Val Gly Tyr  
 1 5 10

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<400> 831

Pro Tyr Gly Gly Ser Gly Phe Gly Ser Tyr  
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<210> 832  
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<400> 832

Gly Ser Gly Tyr Gly Tyr Gly Tyr Gly Tyr  
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<400> 833

Gly Tyr Gly Tyr Gly Gly Tyr Thr Asp Pro Arg  
 1 5 10

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<400> 834

Gly Val Asn Pro Thr Ala Gln Ser Ser Gly  
 1 5 10

<210> 835  
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<400> 835

Ser Leu Tyr Gly Ser Gln Ile Tyr Ala Leu  
 1 5 10

<210> 836  
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<220>  
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<400> 836

Cys Asn Gln Phe Tyr Thr Pro Ala Ala Thr  
 1 5 10

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<220>  
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<400> 837

Gly Leu Tyr Val Asp Gln Tyr Leu Tyr His  
 1 5 10

<210> 838  
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<212> PRT  
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<220>  
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<400> 838

Tyr Cys Val Val Asp Pro Gln Glu  
 1 5

<210> 839  
 <211> 5  
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<220>  
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<400> 839

Gly Val Asn Pro Thr  
 1 5

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<220>  
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<400> 840

Ala Gln Ser Ser Gly Ser Leu Tyr Gly Ser  
 1 5 10

<210> 841  
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<220>  
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<400> 841

Gln Ile Tyr Ala Leu Cys Asn Gln Phe Tyr  
 1 5 10

<210> 842  
 <211> 10  
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<220>  
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<400> 842

Thr Pro Ala Ala Thr Gly Leu Tyr Val Asp  
1 5 10

<210> 843  
<211> 10  
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<220>  
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<400> 843

Gln Tyr Leu Tyr His Tyr Cys Val Val Asp  
1 5 10

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<220>  
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<400> 844

Pro Gln Glu  
1

<210> 845  
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<220>  
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<221> misc\_feature  
<222> (2)..(2)  
<223> Xaa can be any naturally occurring amino acid

<400> 845

Tyr Xaa Arg Phe  
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<210> 846  
<211> 13  
<212> PRT  
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<400> 846

Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln  
 1 5 10

<210> 847  
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<220>  
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<400> 847

Ser Val Thr Val His Ser Ser Glu Pro  
 1 5

<210> 848  
 <211> 10  
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 <213> Artificial Sequence

<220>  
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<400> 848

Lys Val Phe Asp Ser Leu Leu Asn Leu Ser  
 1 5 10

<210> 849  
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 <212> PRT  
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<220>  
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<400> 849

Asp Arg Gly Tyr Gly Thr Ser Leu Leu  
 1 5

<210> 850  
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<400> 850

Gly Tyr Gly Tyr Gly Tyr Gly Tyr Gly  
 1 5

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<220>

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<400> 851

Gly Ser Gly Phe Gly Ser Tyr Gly Ser  
1 5

<210> 852

<211> 9

<212> PRT

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<220>

<223> Synthetic construct

<400> 852

Lys Phe Asp Gln Gly Asp Thr Thr Arg  
1 5

<210> 853

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic construct

<400> 853

Lys Val Tyr Asp Ser Leu Leu Ala Leu Pro  
1 5 10

<210> 854

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic construct

<400> 854

Gly Glu Val Lys Val Lys Leu Ile Val  
1 5

<210> 855

<211> 10

<212> PRT

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<220>

<223> Synthetic construct

<400> 855

Asn Arg Ile Val Gln Glu Phe Tyr Asp Pro  
 1 5 10

<210> 856  
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<400> 856

Val Ser Glu Glu Gly Gly Asn Ser Tyr  
 1 5

<210> 857  
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<220>  
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<400> 857

Leu Val Cys Tyr Asn Asn Lys Ile Thr  
 1 5

<210> 858  
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<220>  
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<400> 858

Ile Val Val Arg Glu Phe Tyr Asp Pro Ser  
 1 5 10

<210> 859  
 <211> 9  
 <212> PRT  
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<220>  
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<400> 859

Tyr Gly Tyr Gly Gly Tyr Thr Asp Pro  
 1 5

<210> 860  
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<220>

<223> Synthetic construct

<400> 860

Val Val Gln Ser Thr Gly His Met Gln Cys  
1 5 10

<210> 861

<211> 10

<212> PRT

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<220>

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<400> 861

Tyr Ala Gly Asp Asn Ile Val Thr Ala Gln  
1 5 10

<210> 862

<211> 10

<212> PRT

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<400> 862

Val Ser Gln Ser Thr Gly Gln Ile Gln Cys  
1 5 10

<210> 863

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<400> 863

Tyr Val Gly Ala Ser Ile Val Thr Ala Val  
1 5 10

<210> 864

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<223> Synthetic construct

<400> 864



Phe Leu Asp His Asn Ile Val Thr Ala Gln  
 1 5 10

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<400> 865

Gly Phe Ser Ser Pro Arg Val Glu Trp  
 1 5

<210> 866  
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<400> 866

Gly Val Asn Pro Thr Ala Gln Ser Ser  
 1 5

<210> 867  
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<400> 867

Gly Ser Leu Tyr Gly Ser Gln Ile Tyr  
 1 5

<210> 868  
 <211> 10  
 <212> PRT  
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<400> 868  
 Ile Gly Ser Asn Ile Ile Thr Ser Gln Asn  
 1 5 10

<210> 869  
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 <212> PRT  
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<400> 869

Val Pro Val Ser Gln Lys Tyr Glu Leu Gly  
 1 5 10

<210> 870  
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<220>  
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<400> 870

Asn Ile Trp Glu Gly Leu Trp Met Asn Cys  
 1 5 10

<210> 871  
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<400> 871

Phe Ile Gly Ser Asn Ile Val Thr Ser Gln  
 1 5 10

<210> 872  
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<400> 872

Val Val Gln Ser Thr Gly Gln Met Gln Cys  
 1 5 10

<210> 873  
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 <212> PRT  
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<220>  
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<400> 873

Phe Ile Gly Ser Asn Ile Ile Thr Ser Gln  
 1 5 10

<210> 874  
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<400> 874

Ala Met Tyr Glu Gly Leu Trp Met Ser Cys  
 1 5 10

<210> 875  
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<400> 875

Gly Gly Ser Val Gly Tyr Pro Tyr Gly  
 1 5

<210> 876  
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<220>  
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<400> 876

Thr Ile Trp Glu Gly Leu Trp Met Asn Cys  
 1 5 10

<210> 877  
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 <212> PRT  
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<220>  
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<400> 877

Asp Ile Tyr Ser Thr Leu Leu Gly Leu Pro  
 1 5 10

<210> 878  
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 <212> PRT  
 <213> Artificial Sequence

&lt;220&gt;

&lt;223&gt; Synthetic construct

&lt;400&gt; 878

Gly	Phe	Ser	Leu	Gly	Leu	Trp	Met	Glu	Cys
1				5					10

&lt;210&gt; 879

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Synthetic construct

&lt;400&gt; 879

Lys	Val	Tyr	Asp	Ser	Val	Leu	Ala	Leu	Ser
1				5					10

&lt;210&gt; 880

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Synthetic construct

&lt;400&gt; 880

Ala	Thr	His	Ser	Thr	Gly	Ile	Thr	Gln	Cys
1				5					10

&lt;210&gt; 881

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Synthetic construct

&lt;400&gt; 881

Thr	Thr	Trp	Leu	Gly	Leu	Trp	Met	Ser	Cys
1				5					10

&lt;210&gt; 882

&lt;211&gt; 5

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Synthetic construct

&lt;400&gt; 882

Val	Leu	Pro	Pro	Ser
1				5

<210> 883  
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 <212> PRT  
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<220>  
 <223> Synthetic construct

<400> 883

Tyr Glu Asp Arg Val Thr Phe  
 1 5

<210> 884  
 <211> 5  
 <212> PRT  
 <213> Artificial Sequence

<220>  
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<400> 884

Pro Arg Val Glu Trp  
 1 5

<210> 885  
 <211> 10  
 <212> PRT  
 <213> Artificial Sequence

<220>  
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<400> 885

Gly Phe Ser Lys Gly Leu Trp Met Glu Cys  
 1 5 10

<210> 886  
 <211> 10  
 <212> PRT  
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<220>  
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<400> 886

Thr Thr Trp Lys Gly Leu Trp Met Ser Cys  
 1 5 10

<210> 887  
 <211> 299  
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&lt;220&gt;

&lt;223&gt; synthetic construct

&lt;400&gt; 887

Met Gly Thr Lys Ala Gln Val Glu Arg Lys Leu Leu Cys Leu Phe Ile  
 1 5 10 15

Leu Ala Ile Leu Leu Cys Ser Leu Ala Leu Gly Ser Val Thr Val His  
 20 25 30

Ser Ser Glu Pro Glu Val Arg Ile Pro Glu Asn Asn Pro Val Lys Leu  
 35 40 45

Ser Cys Ala Tyr Ser Gly Phe Ser Ser Pro Arg Val Glu Trp Lys Phe  
 50 55 60

Asp Gln Gly Asp Thr Thr Arg Leu Val Cys Tyr Asn Asn Lys Ile Thr  
 65 70 75 80

Ala Ser Tyr Glu Asp Arg Val Thr Phe Leu Pro Thr Gly Ile Thr Phe  
 85 90 95

Lys Ser Val Thr Arg Glu Asp Thr Gly Thr Tyr Thr Cys Met Val Ser  
 100 105 110

Glu Glu Gly Gly Asn Ser Tyr Gly Glu Val Lys Val Lys Leu Ile Val  
 115 120 125

Leu Val Pro Pro Ser Lys Pro Thr Val Asn Ile Pro Ser Ser Ala Thr  
 130 135 140

Ile Gly Asn Arg Ala Val Leu Thr Cys Ser Glu Gln Asp Gly Ser Pro  
 145 150 155 160

Pro Ser Glu Tyr Thr Trp Phe Lys Asp Gly Ile Val Met Pro Thr Asn  
 165 170 175

Pro Lys Ser Thr Arg Ala Phe Ser Asn Ser Ser Tyr Val Leu Asn Pro  
 180 185 190

Thr Thr Gly Glu Leu Val Phe Asp Pro Leu Ser Ala Ser Asp Thr Gly  
 195 200 205

Glu Tyr Ser Cys Glu Ala Arg Asn Gly Tyr Gly Thr Pro Met Thr Ser  
 210 215 220

Asn Ala Val Arg Met Glu Ala Val Glu Arg Asn Val Gly Val Ile Val  
 225 230 235 240

Ala Ala Val Leu Val Thr Leu Ile Leu Leu Gly Ile Leu Val Phe Gly  
245 250 255

Ile Trp Phe Ala Tyr Ser Arg Gly His Phe Asp Arg Thr Lys Lys Gly  
260 265 270

Thr Ser Ser Lys Lys Val Ile Tyr Ser Gln Pro Ser Ala Arg Ser Glu  
275 280 285

Gly Glu Phe Lys Gln Thr Ser Ser Phe Leu Val  
290 295

<210> 888

<211> 310

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic construct

<400> 888

Met Ala Leu Arg Arg Pro Pro Arg Leu Arg Leu Cys Ala Arg Leu Pro  
1 5 10 15

Asp Phe Phe Leu Leu Leu Leu Phe Arg Gly Cys Leu Ile Gly Ala Val  
20 25 30

Asn Leu Lys Ser Ser Asn Arg Thr Pro Val Val Gln Glu Phe Glu Ser  
35 40 45

Val Glu Leu Ser Cys Ile Ile Thr Asp Ser Gln Thr Ser Asp Pro Arg  
50 55 60

Ile Glu Trp Lys Lys Ile Gln Asp Glu Gln Thr Thr Tyr Val Phe Phe  
65 70 75 80

Asp Asn Lys Ile Gln Gly Asp Leu Ala Gly Arg Ala Glu Ile Leu Gly  
85 90 95

Lys Thr Ser Leu Lys Ile Trp Asn Val Thr Arg Arg Asp Ser Ala Leu  
100 105 110

Tyr Arg Cys Glu Val Val Ala Arg Asn Asp Arg Lys Glu Ile Asp Glu  
115 120 125

Ile Val Ile Glu Leu Thr Val Gln Val Lys Pro Val Thr Pro Val Cys  
130 135 140

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Arg Val Pro Lys Ala Val Pro Val Gly Lys Met Ala Thr Leu His Cys  
145 150 155 160

Gln Glu Ser Glu Gly His Pro Arg Pro His Tyr Ser Trp Tyr Arg Asn  
165 170 175

Asp Val Pro Leu Pro Thr Asp Ser Arg Ala Asn Pro Arg Phe Arg Asn  
180 185 190

Ser Ser Phe His Leu Asn Ser Glu Thr Gly Thr Leu Val Phe Thr Ala  
195 200 205

Val His Lys Asp Asp Ser Gly Gln Tyr Tyr Cys Ile Ala Ser Asn Asp  
210 215 220

Ala Gly Ser Ala Arg Cys Glu Glu Gln Glu Met Glu Val Tyr Asp Leu  
225 230 235 240

Asn Ile Gly Gly Ile Ile Gly Gly Val Leu Val Val Leu Ala Val Leu  
245 250 255

Ala Leu Ile Thr Leu Gly Ile Cys Cys Ala Tyr Arg Arg Gly Tyr Phe  
260 265 270

Ile Asn Asn Lys Gln Asp Gly Glu Ser Tyr Lys Asn Pro Gly Lys Pro  
275 280 285

Asp Gly Val Asn Tyr Ile Arg Thr Asp Glu Glu Gly Asp Phe Arg His  
290 295 300

Lys Ser Ser Phe Val Ile  
305 310

<210> 889  
<211> 298  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic construct

<400> 889

Met Ala Arg Arg Ser Arg His Arg Leu Leu Leu Leu Leu Arg Tyr  
1 5 10 15

Leu Val Val Ala Leu Gly Tyr His Lys Ala Tyr Gly Phe Ser Ala Pro  
20 25 30



Lys Asp Gln Gln Val Val Thr Ala Val Glu Tyr Gln Glu Ala Ile Leu  
 35 40 45  
 Ala Cys Lys Thr Pro Lys Lys Thr Val Ser Ser Arg Leu Glu Trp Lys  
 50 55 60  
 Lys Leu Gly Arg Ser Val Ser Phe Val Tyr Tyr Gln Gln Thr Leu Gln  
 65 70 75 80  
 Gly Asp Phe Lys Asn Arg Ala Glu Met Ile Asp Phe Asn Ile Arg Ile  
 85 90 95  
 Lys Asn Val Thr Arg Ser Asp Ala Gly Lys Tyr Arg Cys Glu Val Ser  
 100 105 110  
 Ala Pro Ser Glu Gln Gly Gln Asn Leu Glu Glu Asp Thr Val Thr Leu  
 115 120 125  
 Glu Val Leu Val Ala Pro Ala Val Pro Ser Cys Glu Val Pro Ser Ser  
 130 135 140  
 Ala Leu Ser Gly Thr Val Val Glu Leu Arg Cys Gln Asp Lys Glu Gly  
 145 150 155 160  
 Asn Pro Ala Pro Glu Tyr Thr Trp Phe Lys Asp Gly Ile Arg Leu Leu  
 165 170 175  
 Glu Asn Pro Arg Leu Gly Ser Gln Ser Thr Asn Ser Ser Tyr Thr Met  
 180 185 190  
 Asn Thr Lys Thr Gly Thr Leu Gln Phe Asn Thr Val Ser Lys Leu Asp  
 195 200 205  
 Thr Gly Glu Tyr Ser Cys Glu Ala Arg Asn Ser Val Gly Tyr Arg Arg  
 210 215 220  
 Cys Pro Gly Lys Arg Met Gln Val Asp Asp Leu Asn Ile Ser Gly Ile  
 225 230 235 240  
 Ile Ala Ala Val Val Val Val Ala Leu Val Ile Ser Val Cys Gly Leu  
 245 250 255  
 Gly Val Cys Tyr Ala Gln Arg Lys Gly Tyr Phe Ser Lys Glu Thr Ser  
 260 265 270  
 Phe Gln Lys Ser Asn Ser Ser Ser Lys Ala Thr Thr Met Ser Glu Asn  
 275 280 285

Asp Phe Lys His Thr Lys Ser Phe Ile Ile  
 290 295

<210> 890  
 <211> 211  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Synthetic construct

<400> 890

Met Ala Asn Ala Gly Leu Gln Leu Leu Gly Phe Ile Leu Ala Phe Leu  
 1 5 10 15

Gly Trp Ile Gly Ala Ile Val Ser Thr Ala Leu Pro Gln Trp Arg Ile  
 20 25 30

Tyr Ser Tyr Ala Gly Asp Asn Ile Val Thr Ala Gln Ala Met Tyr Glu  
 35 40 45

Gly Leu Trp Met Ser Cys Val Ser Gln Ser Thr Gly Gln Ile Gln Cys  
 50 55 60

Lys Val Phe Asp Ser Leu Leu Asn Leu Ser Ser Thr Leu Gln Ala Thr  
 65 70 75 80

Arg Ala Leu Met Val Val Gly Ile Leu Leu Gly Val Ile Ala Ile Phe  
 85 90 95

Val Ala Thr Val Gly Met Lys Cys Met Lys Cys Leu Glu Asp Asp Glu  
 100 105 110

Val Gln Lys Met Arg Met Ala Val Ile Gly Gly Ala Ile Phe Leu Leu  
 115 120 125

Ala Gly Leu Ala Ile Leu Val Ala Thr Ala Trp Tyr Gly Asn Arg Ile  
 130 135 140

Val Gln Glu Phe Tyr Asp Pro Met Thr Pro Val Asn Ala Arg Tyr Glu  
 145 150 155 160

Phe Gly Gln Ala Leu Phe Thr Gly Trp Ala Ala Ala Ser Leu Cys Leu  
 165 170 175

Leu Gly Gly Ala Leu Leu Cys Cys Ser Cys Pro Arg Lys Thr Thr Ser  
 180 185 190

Tyr Pro Thr Pro Arg Pro Tyr Pro Lys Pro Ala Pro Ser Ser Gly Lys  
 195 200 205

Asp Tyr Val  
 210

<210> 891  
 <211> 229  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> synthetic construct

<400> 891

Met Ala Ser Leu Gly Leu Gln Leu Val Gly Tyr Ile Leu Gly Leu Leu  
 1 5 10 15

Gly Leu Leu Gly Thr Leu Val Ala Met Leu Leu Pro Ser Trp Lys Thr  
 20 25 30

Ser Ser Tyr Val Gly Ala Ser Ile Val Thr Ala Val Gly Phe Ser Lys  
 35 40 45

Gly Leu Trp Met Glu Cys Ala Thr His Ser Thr Gly Ile Thr Gln Cys  
 50 55 60

Asp Ile Tyr Ser Thr Leu Leu Gly Leu Pro Ala Asp Ile Gln Ala Ala  
 65 70 75 80

Gln Ala Met Met Val Thr Ser Ser Ala Ile Ser Ser Leu Ala Cys Ile  
 85 90 95

Ile Ser Val Val Gly Met Arg Cys Thr Val Phe Cys Gln Glu Ser Arg  
 100 105 110

Ala Lys Asp Arg Val Ala Val Ala Gly Gly Val Phe Phe Ile Leu Gly  
 115 120 125

Gly Leu Leu Gly Phe Ile Pro Val Ala Trp Asn Leu His Gly Ile Leu  
 130 135 140

Arg Asp Phe Tyr Ser Pro Leu Val Pro Asp Ser Met Lys Phe Glu Ile  
 145 150 155 160

Gly Glu Ala Leu Tyr Leu Gly Ile Ile Ser Ser Leu Phe Ser Leu Ile  
 165 170 175

Ala Gly Ile Ile Leu Cys Phe Ser Cys Ser Ser Gln Arg Asn Arg Ser  
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Asn Tyr Tyr Asp Ala Tyr Gln Ala Gln Pro Leu Ala Thr Arg Ser Ser  
 195 200 205

Pro Arg Pro Gly Gln Pro Pro Lys Val Lys Ser Glu Phe Asn Ser Tyr  
 210 215 220

Ser Leu Thr Gly Tyr  
 225

<210> 892  
 <211> 220  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> synthetic construct

<400> 892

Met Ser Met Gly Leu Glu Ile Thr Gly Thr Ala Leu Ala Val Leu Gly  
 1 5 10 15

Trp Leu Gly Thr Ile Val Cys Cys Ala Leu Pro Met Trp Arg Val Ser  
 20 25 30

Ala Phe Ile Gly Ser Asn Ile Ile Thr Ser Gln Asn Ile Trp Glu Gly  
 35 40 45

Leu Trp Met Asn Cys Val Val Gln Ser Thr Gly Gln Met Gln Cys Lys  
 50 55 60

Val Tyr Asp Ser Leu Leu Ala Leu Pro Gln Asp Leu Gln Ala Ala Arg  
 65 70 75 80

Ala Leu Ile Val Val Ala Ile Leu Leu Ala Ala Phe Gly Leu Leu Val  
 85 90 95

Ala Leu Val Gly Ala Gln Cys Thr Asn Cys Val Gln Asp Asp Thr Ala  
 100 105 110

Lys Ala Lys Ile Thr Ile Val Ala Gly Val Leu Phe Leu Leu Ala Ala  
 115 120 125

Leu Leu Thr Leu Val Pro Val Ser Trp Ser Ala Asn Thr Ile Ile Arg  
 130 135 140

Asp Phe Tyr Asn Pro Val Val Pro Glu Ala Gln Lys Arg Glu Met Gly  
 145 150 155 160

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Ala Gly Leu Tyr Val Gly Trp Ala Ala Ala Ala Leu Gln Leu Leu Gly  
165 170 175

Gly Ala Leu Leu Cys Cys Ser Cys Pro Pro Arg Glu Lys Lys Tyr Thr  
180 185 190

Ala Thr Lys Val Val Tyr Ser Ala Pro Arg Ser Thr Gly Pro Gly Ala  
195 200 205

Ser Leu Gly Thr Gly Tyr Asp Arg Lys Asp Tyr Val  
210 215 220

<210> 893  
<211> 209  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic construct

<400> 893

Met Ala Ser Met Gly Leu Gln Val Met Gly Ile Ala Leu Ala Val Leu  
1 5 10 15

Gly Trp Leu Ala Val Met Leu Cys Cys Ala Leu Pro Met Trp Arg Val  
20 25 30

Thr Ala Phe Ile Gly Ser Asn Ile Val Thr Ser Gln Thr Ile Trp Glu  
35 40 45

Gly Leu Trp Met Asn Cys Val Val Gln Ser Thr Gly Gln Met Gln Cys  
50 55 60

Lys Val Tyr Asp Ser Leu Leu Ala Leu Pro Gln Asp Leu Gln Ala Ala  
65 70 75 80

Arg Ala Leu Val Ile Ile Ser Ile Ile Val Ala Ala Leu Gly Val Leu  
85 90 95

Leu Ser Val Val Gly Gly Lys Cys Thr Asn Cys Leu Glu Asp Glu Ser  
100 105 110

Ala Lys Ala Lys Thr Met Ile Val Ala Gly Val Val Phe Leu Leu Ala  
115 120 125

Gly Leu Met Val Ile Val Pro Val Ser Trp Thr Ala His Asn Ile Ile  
130 135 140

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Gln Asp Phe Tyr Asn Pro Leu Val Ala Ser Gly Gln Lys Arg Glu Met  
145 150 155 160

Gly Ala Ser Leu Tyr Val Gly Trp Ala Ala Ser Gly Leu Leu Leu Leu  
165 170 175

Gly Gly Gly Leu Leu Cys Cys Asn Cys Pro Pro Arg Thr Asp Lys Pro  
180 185 190

Tyr Ser Ala Lys Tyr Ser Ala Ala Arg Ser Ala Ala Ala Ser Asn Tyr  
195 200 205

Val

<210> 894  
<211> 218  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> synthetic construct

<400> 894

Met Gly Ser Ala Ala Leu Glu Ile Leu Gly Leu Val Leu Cys Leu Val  
1 5 10 15

Gly Trp Gly Gly Leu Ile Leu Ala Cys Gly Leu Pro Met Trp Gln Val  
20 25 30

Thr Ala Phe Leu Asp His Asn Ile Val Thr Ala Gln Thr Thr Trp Lys  
35 40 45

Gly Leu Trp Met Ser Cys Val Val Gln Ser Thr Gly His Met Gln Cys  
50 55 60

Lys Val Tyr Asp Ser Val Leu Ala Leu Ser Thr Glu Val Gln Ala Ala  
65 70 75 80

Arg Ala Leu Thr Val Ser Ala Val Leu Leu Ala Phe Val Ala Leu Phe  
85 90 95

Val Thr Leu Ala Gly Ala Gln Cys Thr Thr Cys Val Ala Pro Gly Pro  
100 105 110

Ala Lys Ala Arg Val Ala Leu Thr Gly Gly Val Leu Tyr Leu Phe Cys  
115 120 125

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Gly Leu Leu Ala Leu Val Pro Leu Cys Trp Phe Ala Asn Ile Val Val  
130 135 140

Arg Glu Phe Tyr Asp Pro Ser Val Pro Val Ser Gln Lys Tyr Glu Leu  
145 150 155 160

Gly Ala Ala Leu Tyr Ile Gly Trp Ala Ala Thr Ala Leu Leu Met Val  
165 170 175

Gly Gly Cys Leu Leu Cys Cys Gly Ala Trp Val Cys Thr Gly Arg Pro  
180 185 190

Asp Leu Ser Phe Pro Val Lys Tyr Ser Ala Pro Arg Arg Pro Thr Ala  
195 200 205

Thr Gly Asp Tyr Asp Lys Lys Asn Tyr Val  
210 215

<210> 895  
<211> 220  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic construct

<400> 895

Met Ala Ser Ala Gly Met Gln Ile Leu Gly Val Val Leu Thr Leu Leu  
1 5 10 15

Gly Trp Val Asn Gly Leu Val Ser Cys Ala Leu Pro Met Trp Lys Val  
20 25 30

Thr Ala Phe Ile Gly Asn Ser Ile Val Val Ala Gln Val Val Trp Glu  
35 40 45

Gly Leu Trp Met Ser Cys Val Val Gln Ser Thr Gly Gln Met Gln Cys  
50 55 60

Lys Val Tyr Asp Ser Leu Leu Ala Leu Pro Gln Asp Leu Gln Ala Ala  
65 70 75 80

Arg Ala Leu Cys Val Ile Ala Leu Leu Val Ala Leu Phe Gly Leu Leu  
85 90 95

Val Tyr Leu Ala Gly Ala Lys Cys Thr Thr Cys Val Glu Glu Lys Asp  
100 105 110

Ser Lys Ala Arg Leu Val Leu Thr Ser Gly Ile Val Phe Val Ile Ser  
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Gly Val Leu Thr Leu Ile Pro Val Cys Trp Thr Ala His Ala Val Ile  
130 135 140

Arg Asp Phe Tyr Asn Pro Leu Val Ala Glu Ala Gln Lys Arg Glu Leu  
145 150 155 160

Gly Ala Ser Leu Tyr Leu Gly Trp Ala Ala Ser Gly Leu Leu Leu Leu  
165 170 175

Gly Gly Gly Leu Leu Cys Cys Thr Cys Pro Ser Gly Gly Ser Gln Gly  
180 185 190

Pro Ser His Tyr Met Ala Arg Tyr Ser Thr Ser Ala Pro Ala Ile Ser  
195 200 205

Arg Gly Pro Ser Glu Tyr Pro Thr Lys Asn Tyr Val  
210 215 220

<210> 896  
<211> 211  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic construct

<400> 896

Met Ala Asn Ser Gly Leu Gln Leu Leu Gly Phe Ser Met Ala Leu Leu  
1 5 10 15

Gly Trp Val Gly Leu Val Ala Cys Thr Ala Ile Pro Gln Trp Gln Met  
20 25 30

Ser Ser Tyr Ala Gly Asp Asn Ile Ile Thr Ala Gln Ala Met Tyr Lys  
35 40 45

Gly Leu Trp Met Asp Cys Val Thr Gln Ser Thr Gly Met Met Ser Cys  
50 55 60

Lys Met Tyr Asp Ser Val Leu Ala Leu Ser Ala Ala Leu Gln Ala Thr  
65 70 75 80

Arg Ala Leu Met Val Val Ser Leu Val Leu Gly Phe Leu Ala Met Phe  
85 90 95

Val Ala Thr Met Gly Met Lys Cys Thr Arg Cys Gly Gly Asp Asp Lys  
100 105 110



Val Lys Lys Ala Arg Ile Ala Met Gly Gly Gly Ile Ile Phe Ile Val  
 115 120 125

Ala Gly Leu Ala Ala Leu Val Ala Cys Ser Trp Tyr Gly His Gln Ile  
 130 135 140

Val Thr Asp Phe Tyr Asn Pro Leu Ile Pro Thr Asn Ile Lys Tyr Glu  
 145 150 155 160

Phe Gly Pro Ala Ile Phe Ile Gly Trp Ala Gly Ser Ala Leu Val Ile  
 165 170 175

Leu Gly Gly Ala Leu Leu Ser Cys Ser Cys Pro Gly Asn Glu Ser Lys  
 180 185 190

Ala Gly Tyr Arg Ala Pro Arg Ser Tyr Pro Lys Ser Asn Ser Ser Lys  
 195 200 205

Glu Tyr Val  
 210

<210> 897  
 <211> 225  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Synthetic construct

<400> 897

Met Ala Thr His Ala Leu Glu Ile Ala Gly Leu Phe Leu Gly Gly Val  
 1 5 10 15

Gly Met Val Gly Thr Val Ala Val Thr Val Met Pro Gln Trp Arg Val  
 20 25 30

Ser Ala Phe Ile Glu Asn Asn Ile Val Val Phe Glu Asn Phe Trp Glu  
 35 40 45

Gly Leu Trp Met Asn Cys Val Arg Gln Ala Asn Ile Arg Met Gln Cys  
 50 55 60

Lys Ile Tyr Asp Ser Leu Leu Ala Leu Ser Pro Asp Leu Gln Ala Ala  
 65 70 75 80

Arg Gly Leu Met Cys Ala Ala Ser Val Met Ser Phe Leu Ala Phe Met  
 85 90 95

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Met Ala Ile Leu Gly Met Lys Cys Thr Arg Cys Thr Gly Asp Asn Glu  
100 105 110

Lys Val Lys Ala His Ile Leu Leu Thr Ala Gly Ile Ile Phe Ile Ile  
115 120 125

Thr Gly Met Val Val Leu Ile Pro Val Ser Trp Val Ala Asn Ala Ile  
130 135 140

Ile Arg Asp Phe Tyr Asn Ser Ile Val Asn Val Ala Gln Lys Arg Glu  
145 150 155 160

Leu Gly Glu Ala Leu Tyr Leu Gly Trp Thr Thr Ala Leu Val Leu Ile  
165 170 175

Val Gly Gly Ala Leu Phe Cys Cys Val Phe Cys Cys Asn Glu Lys Ser  
180 185 190

Ser Ser Tyr Arg Tyr Ser Ile Pro Ser His Arg Thr Thr Gln Lys Ser  
195 200 205

Tyr His Thr Gly Lys Lys Ser Pro Ser Val Tyr Ser Arg Ser Gln Tyr  
210 215 220

Val  
225

<210> 898  
<211> 217  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Synthetic construct

<400> 898

Met Ala Ser Thr Gly Leu Glu Leu Leu Gly Met Thr Leu Ala Val Leu  
1 5 10 15

Gly Trp Leu Gly Thr Leu Val Ser Cys Ala Leu Pro Leu Trp Lys Val  
20 25 30

Thr Ala Phe Ile Gly Asn Ser Ile Val Val Ala Gln Val Val Trp Glu  
35 40 45

Gly Leu Trp Met Ser Cys Val Val Gln Ser Thr Gly Gln Met Gln Cys  
50 55 60

Lys Val Tyr Asp Ser Leu Leu Ala Leu Pro Gln Asp Leu Gln Ala Ala  
65 70 75 80

Arg Ala Leu Cys Val Ile Ala Leu Leu Leu Ala Leu Leu Gly Leu Leu  
85 90 95

Val Ala Ile Thr Gly Ala Gln Cys Thr Thr Cys Val Glu Asp Glu Gly  
100 105 110

Ala Lys Ala Arg Ile Val Leu Thr Ala Gly Val Ile Leu Leu Leu Ala  
115 120 125

Gly Ile Leu Val Leu Ile Pro Val Cys Trp Thr Ala His Ala Ile Ile  
130 135 140

Gln Asp Phe Tyr Asn Pro Leu Val Ala Glu Ala Leu Lys Arg Glu Leu  
145 150 155 160

Gly Ala Ser Leu Tyr Leu Gly Trp Ala Ala Ala Leu Leu Met Leu  
165 170 175

Gly Gly Gly Leu Leu Cys Cys Thr Cys Pro Pro Pro Gln Val Glu Arg  
180 185 190

Pro Arg Gly Pro Arg Leu Gly Tyr Ser Ile Pro Ser Arg Ser Gly Ala  
195 200 205

Ser Gly Leu Asp Lys Arg Asp Tyr Val  
210 215

<210> 899  
<211> 228  
<212> PRT  
<213> Artificial sequence

<220>  
<223> synthetic construct

<400> 899

Met Ala Ser Thr Ala Ser Glu Ile Ile Ala Phe Met Val Ser Ile Ser  
1 5 10 15

Gly Trp Val Leu Val Ser Ser Thr Leu Pro Thr Asp Tyr Trp Lys Val  
20 25 30

Ser Thr Ile Asp Gly Thr Val Ile Thr Thr Ala Thr Tyr Trp Ala Asn  
35 40 45

Leu Trp Lys Ala Cys Val Thr Asp Ser Thr Gly Val Ser Asn Cys Lys  
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Asp Phe Pro Ser Met Leu Ala Leu Asp Gly Tyr Ile Gln Ala Cys Arg  
65 70 75 80

Gly Leu Met Ile Ala Ala Val Ser Leu Gly Phe Phe Gly Ser Ile Phe  
85 90 95

Ala Leu Phe Gly Met Lys Cys Thr Lys Val Gly Gly Ser Asp Lys Ala  
100 105 110

Lys Ala Lys Ile Ala Cys Leu Ala Gly Ile Val Phe Ile Leu Ser Gly  
115 120 125

Leu Cys Ser Met Thr Gly Cys Ser Leu Tyr Ala Asn Lys Ile Thr Thr  
130 135 140

Glu Phe Phe Asp Pro Leu Phe Val Glu Gln Lys Tyr Glu Leu Gly Ala  
145 150 155 160

Ala Leu Phe Ile Gly Trp Ala Gly Ala Ser Leu Cys Ile Ile Gly Gly  
165 170 175

Val Ile Phe Cys Phe Ser Ile Ser Asp Asn Asn Lys Thr Pro Arg Tyr  
180 185 190

Thr Tyr Asn Gly Ala Thr Ser Val Met Ser Ser Arg Thr Lys Tyr His  
195 200 205

Gly Gly Glu Asp Phe Lys Thr Thr Asn Pro Ser Lys Gln Phe Asp Lys  
210 215 220

Asn Ala Tyr Val  
225

<210> 900  
<211> 522  
<212> PRT  
<213> Artificial sequence

<220>  
<223> Synthetic construct

<400> 900

Met Ser Ser Arg Pro Leu Glu Ser Pro Pro Pro Tyr Arg Pro Asp Glu  
1 5 10 15

Phe Lys Pro Asn His Tyr Ala Pro Ser Asn Asp Ile Tyr Gly Gly Glu  
20 25 30

Met His Val Arg Pro Met Leu Ser Gln Pro Ala Tyr Ser Phe Tyr Pro  
 35 40 45  
 Glu Asp Glu Ile Leu His Phe Tyr Lys Trp Thr Ser Pro Pro Gly Val  
 50 55 60  
 Ile Arg Ile Leu Ser Met Leu Ile Ile Val Met Cys Ile Ala Ile Phe  
 65 70 75 80  
 Ala Cys Val Ala Ser Thr Leu Ala Trp Asp Arg Gly Tyr Gly Thr Ser  
 85 90 95  
 Leu Leu Gly Gly Ser Val Gly Tyr Pro Tyr Gly Gly Ser Gly Phe Gly  
 100 105 110  
 Ser Tyr Gly Ser Gly Tyr Gly Tyr Gly Tyr Gly Tyr Gly Tyr  
 115 120 125  
 Gly Gly Tyr Thr Asp Pro Arg Ala Ala Lys Gly Phe Met Leu Ala Met  
 130 135 140  
 Ala Ala Phe Cys Phe Ile Ala Ala Leu Val Ile Phe Val Thr Ser Val  
 145 150 155 160  
 Ile Arg Ser Glu Met Ser Arg Thr Arg Arg Tyr Tyr Leu Ser Val Ile  
 165 170 175  
 Ile Val Ser Ala Ile Leu Gly Ile Met Val Phe Ile Ala Thr Ile Val  
 180 185 190  
 Tyr Ile Met Gly Val Asn Pro Thr Ala Gln Ser Ser Gly Ser Leu Tyr  
 195 200 205  
 Gly Ser Gln Ile Tyr Ala Leu Cys Asn Gln Phe Tyr Thr Pro Ala Ala  
 210 215 220  
 Thr Gly Leu Tyr Val Asp Gln Tyr Leu Tyr His Tyr Cys Val Val Asp  
 225 230 235 240  
 Pro Gln Glu Ala Ile Ala Ile Val Leu Gly Phe Met Ile Ile Val Ala  
 245 250 255  
 Phe Ala Leu Ile Ile Phe Phe Ala Val Lys Thr Arg Arg Lys Met Asp  
 260 265 270  
 Arg Tyr Asp Lys Ser Asn Ile Leu Trp Asp Lys Glu His Ile Tyr Asp  
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Glu Gln Pro Pro Asn Val Glu Glu Trp Val Lys Asn Val Ser Ala Gly  
 290 295 300  
 Thr Gln Asp Val Pro Ser Pro Pro Ser Asp Tyr Val Glu Arg Val Asp  
 305 310 315 320  
 Ser Pro Met Ala Tyr Ser Ser Asn Gly Lys Val Asn Asp Lys Arg Phe  
 325 330 335  
 Tyr Pro Glu Ser Ser Tyr Lys Ser Thr Pro Val Pro Glu Val Val Gln  
 340 345 350  
 Glu Leu Pro Leu Thr Ser Pro Val Asp Asp Phe Arg Gln Pro Arg Tyr  
 355 360 365  
 Ser Ser Gly Gly Asn Phe Glu Thr Pro Ser Lys Arg Ala Pro Ala Lys  
 370 375 380  
 Gly Arg Ala Gly Arg Ser Lys Arg Thr Glu Gln Asp His Tyr Glu Thr  
 385 390 395 400  
 Asp Tyr Thr Thr Gly Gly Glu Ser Cys Asp Glu Leu Glu Glu Asp Trp  
 405 410 415  
 Ile Arg Glu Tyr Pro Pro Ile Thr Ser Asp Gln Gln Arg Gln Leu Tyr  
 420 425 430  
 Lys Arg Asn Phe Asp Thr Gly Leu Gln Glu Tyr Lys Ser Leu Gln Ser  
 435 440 445  
 Glu Leu Asp Glu Ile Asn Lys Glu Leu Ser Arg Leu Asp Lys Glu Leu  
 450 455 460  
 Asp Asp Tyr Arg Glu Glu Ser Glu Glu Tyr Met Ala Ala Ala Asp Glu  
 465 470 475 480  
 Tyr Asn Arg Leu Lys Gln Val Lys Gly Ser Ala Asp Tyr Lys Ser Lys  
 485 490 495  
 Lys Asn His Cys Lys Gln Leu Lys Ser Lys Leu Ser His Ile Lys Lys  
 500 505 510  
 Met Val Gly Asp Tyr Asp Arg Gln Lys Thr  
 515 520